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SDA F0 no: 2

8-15-95

GenCore version 4.5  
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OM protein - protein search, using sw model

Run on: November 16, 2001, 15:44:13 ; Search time 267.62 Seconds  
(without alignments)  
17.638 Million cell updates/sec

Title: US-09-011-797-2  
Perfect score: 86  
Sequence: 1 FGGFTGARKSARKLANQ 17

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 3148936 seqs, 277657034 residues

Total number of hits satisfying chosen parameters: 3148936

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Pending Patents\_AA Main:\*

- 1: /cgn2\_6/ptodata/2/paa/PCTUS\_COMB.pep:\*
- 2: /cgn2\_6/ptodata/2/paa/US06\_COMB.pep:\*
- 3: /cgn2\_6/ptodata/2/paa/US07\_COMB.pep:\*
- 4: /cgn2\_6/ptodata/2/paa/US080\_COMB.pep:\*
- 5: /cgn2\_6/ptodata/2/paa/US081\_COMB.pep:\*
- 6: /cgn2\_6/ptodata/2/paa/US082\_COMB.pep:\*
- 7: /cgn2\_6/ptodata/2/paa/US083\_COMB.pep:\*
- 8: /cgn2\_6/ptodata/2/paa/US084\_COMB.pep:\*
- 9: /cgn2\_6/ptodata/2/paa/US085\_COMB.pep:\*
- 10: /cgn2\_6/ptodata/2/paa/US086\_COMB.pep:\*
- 11: /cgn2\_6/ptodata/2/paa/US087\_COMB.pep:\*
- 12: /cgn2\_6/ptodata/2/paa/US088\_COMB.pep:\*
- 13: /cgn2\_6/ptodata/2/paa/US089\_COMB.pep:\*
- 14: /cgn2\_6/ptodata/2/paa/US090\_COMB.pep:\*
- 15: /cgn2\_6/ptodata/2/paa/US091\_COMB.pep:\*
- 16: /cgn2\_6/ptodata/2/paa/US092\_COMB.pep:\*
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- 21: /cgn2\_6/ptodata/2/paa/US097\_COMB.pep:\*
- 22: /cgn2\_6/ptodata/2/paa/US098\_COMB.pep:\*
- 23: /cgn2\_6/ptodata/2/paa/US099\_COMB.pep:\*
- 24: /cgn2\_6/ptodata/2/paa/US60\_COMB.pep:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,

and is derived by analysis of the total score distribution.

# SUMMARIES

Result		%					
No.	Score	Query	Match	Length	DB	ID	Description
1	86	100.0	17	9	US-08-514-451-5	Sequence 5, Appli	
2	86	100.0	17	9	US-08-553-058A-5	Sequence 5, Appli	
3	86	100.0	17	12	US-08-868-355-1	Sequence 1, Appli	
4	86	100.0	17	13	US-08-927-328-1	Sequence 1, Appli	
5	86	100.0	17	14	US-09-011-797-2	Sequence 2, Appli	
6	86	100.0	17	14	US-09-048-916-5	Sequence 5, Appli	
7	86	100.0	17	14	US-09-048-916-5	Sequence 5, Appli	
8	86	100.0	17	14	US-09-048-916B-5	Sequence 5, Appli	
9	86	100.0	17	15	US-09-114-620-1	Sequence 1, Appli	
10	86	100.0	17	15	US-09-170-919-5	Sequence 5, Appli	
11	86	100.0	17	17	US-09-341-590-39	Sequence 39, Appl	
12	86	100.0	17	20	US-09-657-276-919	Sequence 919, App	
13	86	100.0	17	20	US-09-657-276-926	Sequence 926, App	
14	86	100.0	134	24	US-60-160-202-4306	Sequence 4306, Ap	
15	86	100.0	134	24	US-60-160-203-6127	Sequence 6127, Ap	
16	86	100.0	139	24	US-60-160-203-5358	Sequence 5358, Ap	
17	86	100.0	155	24	US-60-160-202-3958	Sequence 3958, Ap	
18	86	100.0	188	1	PCT-US01-18569-3681	Sequence 3681, Ap	
19	83	96.5	17	13	US-08-927-328-4	Sequence 4, Appli	
20	83	96.5	17	13	US-08-927-328-6	Sequence 6, Appli	
21	83	96.5	17	13	US-08-927-328-8	Sequence 8, Appli	
22	81	94.2	17	9	US-08-514-451-6	Sequence 6, Appli	
23	81	94.2	17	9	US-08-553-058A-6	Sequence 6, Appli	
24	81	94.2	17	14	US-09-048-916-6	Sequence 6, Appli	
25	81	94.2	17	14	US-09-048-916-6	Sequence 6, Appli	
26	81	94.2	17	14	US-09-048-916B-6	Sequence 6, Appli	
27	81	94.2	17	15	US-09-114-620-4	Sequence 4, Appli	
28	81	94.2	17	15	US-09-170-919-6	Sequence 6, Appli	
29	80	93.0	16	13	US-08-927-328-3	Sequence 3, Appli	
30	80	93.0	17	13	US-08-927-328-7	Sequence 7, Appli	
31	74	86.0	17	13	US-08-927-328-5	Sequence 5, Appli	
32	68	79.1	15	13	US-08-927-328-20	Sequence 20, Appl	
33	68	79.1	16	13	US-08-927-328-17	Sequence 17, Appl	
34	64	74.4	17	13	US-08-927-328-13	Sequence 13, Appl	
35	63	73.3	14	13	US-08-927-328-30	Sequence 30, Appl	
36	62	72.1	15	13	US-08-927-328-12	Sequence 12, Appl	
37	62	72.1	16	13	US-08-927-328-11	Sequence 11, Appl	
38	62	72.1	16	13	US-08-927-328-15	Sequence 15, Appl	
39	61	70.9	16	13	US-08-927-328-19	Sequence 19, Appl	
40	57	66.3	15	13	US-08-927-328-9	Sequence 9, Appli	
41	57	66.3	15	13	US-08-927-328-14	Sequence 14, Appl	
42	57	66.3	16	13	US-08-927-328-10	Sequence 10, Appl	
43	57	66.3	16	13	US-08-927-328-16	Sequence 16, Appl	
44	57	66.3	16	13	US-08-927-328-18	Sequence 18, Appl	
45	44.5	51.7	319	15	US-09-198-452A-173	Sequence 173, App	

# ALIGNMENTS

RESULT 1  
 US-08-514-451-5  
 ; Sequence 5, Application US/08514451  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Bunzow, James R  
 ; APPLICANT: Grandy, David K  
 ; APPLICANT: Civelli, Olivier  
 ; APPLICANT: Reinscheid, Rainer K  
 ; APPLICANT: Nothacker, Hans-Peter  
 ; APPLICANT: Monsma, Frederick J  
 ; TITLE OF INVENTION: A Novel Mammalian Methadone-Specific  
 ; TITLE OF INVENTION: Opioid Receptor Gene and Uses  
 ; NUMBER OF SEQUENCES: 6  
 ; CORRESPONDENCE ADDRESS:  
 ; ADDRESSEE: Banner & Allegretti, Ltd.  
 ; STREET: 10 South Wacker Drive, Suite 3000  
 ; CITY: Chicago  
 ; STATE: Illinois  
 ; COUNTRY: USA  
 ; ZIP: 60606  
 ; COMPUTER READABLE FORM:  
 ; MEDIUM TYPE: Floppy disk  
 ; COMPUTER: IBM PC compatible  
 ; OPERATING SYSTEM: PC-DOS/MS-DOS  
 ; SOFTWARE: PatentIn Release #1.0, Version #1.25  
 ; CURRENT APPLICATION DATA:  
 ; APPLICATION NUMBER: US/08/514,451  
 ; FILING DATE: 11-AUG-1995  
 ; CLASSIFICATION: 530  
 ; ATTORNEY/AGENT INFORMATION:  
 ; NAME: Noonan, Kevin E  
 ; REGISTRATION NUMBER: 35,303  
 ; REFERENCE/DOCKET NUMBER: 93,311-A  
 ; TELECOMMUNICATION INFORMATION:  
 ; TELEPHONE: 312-715-1000  
 ; TELEFAX: 312-715-1234  
 ; TELEX: 910-221-5317  
 ; INFORMATION FOR SEQ ID NO: 5:  
 ; SEQUENCE CHARACTERISTICS:  
 ; LENGTH: 17 amino acids  
 ; TYPE: amino acid  
 ; TOPOLOGY: linear  
 ; MOLECULE TYPE: peptide  
 US-08-514-451-5

Query Match 100.0%; Score 86; DB 9; Length 17;  
 Best Local Similarity 100.0%; Pred. No. 7.5e-07;  
 Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FGGFTGARKSARKLANQ 17  
 |||||  
 Db 1 FGGFTGARKSARKLANQ 17

RESULT 2  
 US-08-553-058A-5

```

; Sequence 5, Application US/08553058A
; GENERAL INFORMATION:
;   APPLICANT:  Grisel, Judith E.
;   APPLICANT:  Grandy, David K.
;   APPLICANT:  Mogil, Jeffrey S.
;   TITLE OF INVENTION:  Opioid Antagonists and Methods of Their Use
;   NUMBER OF SEQUENCES:  10
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE:  Klarquist Sparkman Campbell Leigh
;     ADDRESSEE:  & Whinston LLP
;     STREET:    121 S.W. Salmon, Suite 1600
;     CITY:      Portland
;     STATE:     Oregon
;     COUNTRY:   USA
;     ZIP:       97204
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE:  Floppy Disk
;     COMPUTER:     IBM PC compatible
;     OPERATING SYSTEM:  PC-DOS/MS-DOS
;     SOFTWARE:     Patent In Release #1.0, Version WP5.1 ASCII text
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER:  US/08/553,058A
;     FILING DATE:       11/13/95
;     CLASSIFICATION:    514
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:
;     FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;     NAME:  William D. Noonan, M.D.
;     REGISTRATION NUMBER:  30,878
;     REFERENCE/DOCKET NUMBER:  899-40006/WDN
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE:  (503) 226-7391
;     TELEFAX:   (503) 228-9446
;   INFORMATION FOR SEQ ID NO:  5:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  17 amino acids
;       TYPE:    amino acid
;       TOPOLOGY:  linear
;     MOLECULE TYPE:
;       DESCRIPTION:  peptide
US-08-553-058A-5

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Query Match          100.0%;  Score 86;  DB 9;  Length 17;
Best Local Similarity 100.0%;  Pred. No. 7.5e-07;
Matches   17;  Conservative   0;  Mismatches   0;  Indels   0;  Gaps   0;

```

```

Qy      1 FGGFTGARKSARKLANQ 17
        ||||||||||||||||
Db      1 FGGFTGARKSARKLANQ 17

```

```

RESULT    3
US-08-868-355-1
; Sequence 1, Application US/08868355
; GENERAL INFORMATION:

```

```

; APPLICANT: Civelli, Olivier
; APPLICANT: Martin, James R.
; APPLICANT: Monsma, Frederick
; APPLICANT: Moreau, Jean-Luc
; APPLICANT: Nothacker, Hans-Peter
; APPLICANT: Reinscheid, Rainer
; TITLE OF INVENTION: MODULATION OF LC132 (OPIOID-LIKE)
; TITLE OF INVENTION: RECEPTOR FUNCTION
; NUMBER OF SEQUENCES: 1
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Lewis J. Kreisler
; STREET: 340 Kingsland Street
; CITY: Nutley
; STATE: New Jersey
; COUNTRY: U.S.A.
; ZIP: 07110-1199
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/868,355
; FILING DATE:
; CLASSIFICATION: -
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: EP 96109462.0
; FILING DATE: 13-JUN-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Kreisler, Lewis J.
; REGISTRATION NUMBER: 38,522
; REFERENCE/DOCKET NUMBER: RAN 4108/361
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (201) 235-4387
; TELEFAX: (201) 235-2363
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 17 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
US-08-868-355-1

```

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Query Match          100.0%;  Score 86;  DB 12;  Length 17;
Best Local Similarity 100.0%;  Pred. No. 7.5e-07;
Matches 17;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

```

```

Qy      1 FGGFTGARKSARKLANQ 17
        |||||
Db      1 FGGFTGARKSARKLANQ 17

```

```

RESULT 4
US-08-927-328-1
; Sequence 1, Application US/08927328

```

```

; GENERAL INFORMATION:
;   APPLICANT:  HOWARD LIPPTON
;   TITLE OF INVENTION:  DIURETIC AND ANTINATRIURETIC RESPONSES
;   TITLE OF INVENTION:  PRODUCED BY ANALOGS OF NOCICEPTIN
;   NUMBER OF SEQUENCES:  31
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE:  Pravel, Hewitt, Kimball & Krieger
;     STREET:  1177 West Loop South, 10th Floor
;     CITY:  Houston
;     STATE:  TX
;     COUNTRY:  USA
;     ZIP:  77027-9095
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE:  Floppy disk
;     COMPUTER:  IBM PC compatible
;     OPERATING SYSTEM:  PC-DOS/MS-DOS
;     SOFTWARE:  PatentIn Release #1.0, Version #1.25
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER:  US/08/927,328
;     FILING DATE:  September 11, 1997
;     CLASSIFICATION:  514
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:
;     FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;     NAME:  Krieger, Paul E.
;     REGISTRATION NUMBER:  25,886
;     REFERENCE/DOCKET NUMBER:  42740/1
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE:  713-850-0909
;     TELEFAX:  713-850-0165
;   INFORMATION FOR SEQ ID NO:  1:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  17 amino acids
;       TYPE:  amino acid
;       STRANDEDNESS:  SINGLE
;       TOPOLOGY:  linear
;     MOLECULE TYPE:  Peptide
US-08-927-328-1

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```

Query Match          100.0%;  Score 86;  DB 13;  Length 17;
Best Local Similarity 100.0%;  Pred. No. 7.5e-07;
Matches  17;  Conservative  0;  Mismatches  0;  Indels  0;  Gaps  0;

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```

Qy      1 FGGFTGARKSARKLANQ 17
        |||||||||||||
Db      1 FGGFTGARKSARKLANQ 17

```

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RESULT  5
US-09-011-797-2
; Sequence 2, Application US/09011797
; GENERAL INFORMATION:
;   APPLICANT:  PARMENTIER, MARC
;   APPLICANT:  VASSART, GILBERT
;   APPLICANT:  MEUNIER, JEAN-CLAUDE

```

```

; APPLICANT: MOLLEREAU, CATHERINE
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING PEPTIDES
; TITLE OF INVENTION: HAVING PRONOCICEPTIVE PROPERTIES
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Knobbe, Martens, Olson & Bear
; STREET: 620 Newport Center Drive 16th Floor
; CITY: Newport Beach
; STATE: CA
; COUNTRY: U.S.A.
; ZIP: 92660
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/011,797
; FILING DATE:
; CLASSIFICATION: 800
; ATTORNEY/AGENT INFORMATION:
; NAME: Altman, Daniel E
; REGISTRATION NUMBER: 34,115
; REFERENCE/DOCKET NUMBER: VANMA72.001APC
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 714-760-0404
; TELEFAX: 714-760-9502
; TELEX:
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 17 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-09-011-797-2

```

```

Query Match          100.0%; Score 86; DB 14; Length 17;
Best Local Similarity 100.0%; Pred. No. 7.5e-07;
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

Qy      1 FGGFTGARKSARKLANQ 17
        |||||||||||||
Db      1 FGGFTGARKSARKLANQ 17

```

```

RESULT 6
US-09-048-916-5
; Sequence 5, Application US/09048916
; GENERAL INFORMATION:
; APPLICANT: Grisel, Judith E.
; APPLICANT: Grandy, David K.
; APPLICANT: Mogil, Jeffrey S.
; APPLICANT: Bunzow, James R.
; APPLICANT: Civelli, Olivier
; APPLICANT: Reinscheid, Rainer Klaus

```



```

; APPLICANT: Nothacker, Hans-Peter
; APPLICANT: Monsma, Frederick James
; TITLE OF INVENTION: Opioid Antagonists and Methods of
; TITLE OF INVENTION: Their Use
; NUMBER OF SEQUENCES: 10
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Klarquist Sparkman Campbell Leigh
; ADDRESSEE: & Whinston LLP
; STREET: 121 S.W. Salmon, Suite 1600
; CITY: Portland
; STATE: Oregon
; COUNTRY: USA
; ZIP: 97204
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy Disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version
; SOFTWARE: WP5.1 ASCII text
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/048,916
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/533,058
; FILING DATE: 3 November 1995
; APPLICATION NUMBER: 08/514,541
; FILING DATE: 11 August 1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Richard J. Polley, Esq.
; REGISTRATION NUMBER: 28,107
; REFERENCE/DOCKET NUMBER: 899-45532/RJP
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (503) 226-7391
; TELEFAX: (503) 228-9446
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 17 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE:
; DESCRIPTION: peptide
US-09-048-916-5

```

```

Query Match          100.0%;  Score 86;  DB 14;  Length 17;
Best Local Similarity 100.0%;  Pred. No. 7.5e-07;
Matches 17;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

```

```

Qy      1 FGGFTGARKSARKLANQ 17
        |||||
Db      1 FGGFTGARKSARKLANQ 17

```

```

RESULT 7
US-09-048-916-5
; Sequence 5, Application US/09048916A

```

```
; GENERAL INFORMATION:
; APPLICANT: Grandy et al.
; TITLE OF INVENTION: Novel mammalian Opioid receptor ligand and uses
; FILE REFERENCE: 49888
; CURRENT APPLICATION NUMBER: US/09/048,916A
; CURRENT FILING DATE: 1999-03-26
; EARLIER APPLICATION NUMBER: 08/514,451
; EARLIER FILING DATE: 1995-08-11
; EARLIER APPLICATION NUMBER: 08/149,093
; EARLIER FILING DATE: 1993-11-08
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 5
;   LENGTH: 17
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence: peptide ligand
US-09-048-916-5
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```
Query Match          100.0%;  Score 86;  DB 14;  Length 17;
Best Local Similarity 100.0%;  Pred. No. 7.5e-07;
Matches 17;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;
```

```
Qy      1 FGGFTGARKSARKLANQ 17
          |||||
Db      1 FGGFTGARKSARKLANQ 17
```

```
RESULT 8
US-09-048-916B-5
; Sequence 5, Application US/09048916B
; GENERAL INFORMATION:
; APPLICANT: Grandy et al.
; TITLE OF INVENTION: Method of screening a compound for binding to MSOR
; FILE REFERENCE: 49888
; CURRENT APPLICATION NUMBER: US/09/048,916B
; CURRENT FILING DATE: 1998-03-26
; PRIOR APPLICATION NUMBER: 08/514,451
; PRIOR FILING DATE: 1995-08-11
; PRIOR APPLICATION NUMBER: 08/149,093
; PRIOR FILING DATE: 1993-11-08
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
;   LENGTH: 17
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence: peptide ligand
US-09-048-916B-5
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Query Match          100.0%;  Score 86;  DB 14;  Length 17;
Best Local Similarity 100.0%;  Pred. No. 7.5e-07;
Matches 17;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;
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Qy 1 FGGFTGARKSARKLANQ 17  
          |||||  
Db 1 FGGFTGARKSARKLANQ 17

RESULT 9

US-09-114-620-1

; Sequence 1, Application US/09114620  
; GENERAL INFORMATION:  
; APPLICANT: Fink-Jensen, Anders  
; APPLICANT: Olsen, Uffe Bang  
; TITLE OF INVENTION: Use of Nociceptin And Nociceptin  
; TITLE OF INVENTION: Analogues For The Manufacture Of A Pharmaceutical  
; TITLE OF INVENTION: Composition For The Treatment Of Hot Flushes  
; FILE REFERENCE: 5286.200-US  
; CURRENT APPLICATION NUMBER: US/09/114,620  
; CURRENT FILING DATE: 1998-07-13  
; EARLIER APPLICATION NUMBER: 60/052,810  
; EARLIER FILING DATE: 1997-07-17  
; EARLIER APPLICATION NUMBER: 0866/97  
; EARLIER FILING DATE: 1997-07-15  
; NUMBER OF SEQ ID NOS: 8  
; SOFTWARE: FastSEQ for Windows Version 3.0  
; SEQ ID NO 1  
; LENGTH: 17  
; TYPE: PRT  
; ORGANISM: Human  
US-09-114-620-1

Query Match 100.0%; Score 86; DB 15; Length 17;  
Best Local Similarity 100.0%; Pred. No. 7.5e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FGGFTGARKSARKLANQ 17  
          |||||  
Db 1 FGGFTGARKSARKLANQ 17

RESULT 10

US-09-170-919-5

; Sequence 5, Application US/09170919  
; GENERAL INFORMATION:  
; APPLICANT: Grisel, Judith E.  
; APPLICANT: Grandy, David K.  
; APPLICANT: Mogil, Jeffrey S.  
; TITLE OF INVENTION: Opioid Antagonists and Methods  
; TITLE OF INVENTION: of Their Use  
; NUMBER OF SEQUENCES: 11  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Klarquist Sparkman Campbell  
; ADDRESSEE: Leigh & Winston LLP  
; STREET: 121 S.W. Salmon, Suite 1600  
; CITY: Portland  
; STATE: Oregon  
; COUNTRY: USA

```

; ZIP: 97204
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy Disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Version WP6, ASCII text
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/170,919
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/553,058
; FILING DATE: 11/3/95
; ATTORNEY/AGENT INFORMATION:
; NAME: William D. Noonan, M.D.
; REGISTRATION NUMBER: 30,878
; REFERENCE/DOCKET NUMBER: 899-40006/WDN
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (503) 226-7391
; TELEFAX: (503) 228-9446
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 17 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE:
; DESCRIPTION: peptide
US-09-170-919-5

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Query Match          100.0%; Score 86; DB 15; Length 17;
Best Local Similarity 100.0%; Pred. No. 7.5e-07;
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy      1 FGGFTGARKSARKLANQ 17
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Db      1 FGGFTGARKSARKLANQ 17

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RESULT 11
US-09-341-590-39
; Sequence 39, Application US/09341590
; GENERAL INFORMATION:
; APPLICANT: Larsen, Bjarne Due
; TITLE OF INVENTION: PHARMACOLOGICALLY ACTIVE PEPTIDE CONJUGATES HAVING A
; TITLE OF INVENTION: REDUCED TENDENCY TOWARDS ENZYMATIC HYDROLYSIS
; FILE REFERENCE: PPT-20479-US
; CURRENT APPLICATION NUMBER: US/09/341,590
; CURRENT FILING DATE: 1999-07-03
; PRIOR APPLICATION NUMBER: DK 0317/98
; PRIOR FILING DATE: 1998-03-09
; NUMBER OF SEQ ID NOS: 121
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 39
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Homo sapiens

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; FEATURE:  
; OTHER INFORMATION: nociceptin  
US-09-341-590-39

Query Match 100.0%; Score 86; DB 17; Length 17;  
Best Local Similarity 100.0%; Pred. No. 7.5e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FGGFTGARKSARKLANQ 17  
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Db 1 FGGFTGARKSARKLANQ 17

RESULT 12

US-09-657-276-919  
; Sequence 919, Application US/09657276  
; GENERAL INFORMATION:  
; APPLICANT: Conjuchem, Inc.  
; APPLICANT: Bridon, Dominique  
; APPLICANT: Ezrin, Alan  
; APPLICANT: Milner, Peter  
; APPLICANT: Holmes, Darren  
; APPLICANT: Thibaudeau, Karen  
; TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM  
; TITLE OF INVENTION: PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD  
; TITLE OF INVENTION: COMPONENTS  
; FILE REFERENCE: 2110  
; CURRENT APPLICATION NUMBER: US/09/657,276  
; CURRENT FILING DATE: 2000-09-07  
; PRIOR APPLICATION NUMBER: 60/134,406  
; PRIOR FILING DATE: 1999-05-17  
; PRIOR APPLICATION NUMBER: 60/153,406  
; PRIOR FILING DATE: 1999-09-10  
; PRIOR APPLICATION NUMBER: 60/159,783  
; PRIOR FILING DATE: 1999-10-18  
; NUMBER OF SEQ ID NOS: 1617  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 919  
; LENGTH: 17  
; TYPE: PRT  
; ORGANISM: Artificial Sequence  
; FEATURE:  
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
; OTHER INFORMATION: Peptide  
US-09-657-276-919

Query Match 100.0%; Score 86; DB 20; Length 17;  
Best Local Similarity 100.0%; Pred. No. 7.5e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FGGFTGARKSARKLANQ 17  
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Db 1 FGGFTGARKSARKLANQ 17

RESULT 13

US-09-657-276-926

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; Sequence 926, Application US/09657276
; GENERAL INFORMATION:
; APPLICANT: Conjuchem, Inc.
; APPLICANT: Bridon, Dominique
; APPLICANT: Ezrin, Alan
; APPLICANT: Milner, Peter
; APPLICANT: Holmes, Darren
; APPLICANT: Thibaudeau, Karen
; TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
; TITLE OF INVENTION: PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD
; TITLE OF INVENTION: COMPONENTS
; FILE REFERENCE: 2110
; CURRENT APPLICATION NUMBER: US/09/657,276
; CURRENT FILING DATE: 2000-09-07
; PRIOR APPLICATION NUMBER: 60/134,406
; PRIOR FILING DATE: 1999-05-17
; PRIOR APPLICATION NUMBER: 60/153,406
; PRIOR FILING DATE: 1999-09-10
; PRIOR APPLICATION NUMBER: 60/159,783
; PRIOR FILING DATE: 1999-10-18
; NUMBER OF SEQ ID NOS: 1617
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 926
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: Peptide
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US-09-657-276-926

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Query Match          100.0%; Score 86; DB 20; Length 17;
Best Local Similarity 100.0%; Pred. No. 7.5e-07;
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy      1 FGGFTGARKSARKLANQ 17
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Db      1 FGGFTGARKSARKLANQ 17
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RESULT 14

US-60-160-202-4306

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; Sequence 4306, Application US/60160202
; GENERAL INFORMATION:
; APPLICANT: BONAZZI, VIVIEN
; TITLE OF INVENTION: ISOLATED HUMAN GPCR PROTEIN, NUCLEIC
; TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN GPCR PROTEINS AND USES
THEREOF
; FILE REFERENCE: CL000114
; CURRENT APPLICATION NUMBER: US/60/160,202
; CURRENT FILING DATE: 1999-10-19
; NUMBER OF SEQ ID NOS: 4392
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 4306
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; LENGTH: 134  
; TYPE: PRT  
; ORGANISM: HUMAN  
US-60-160-202-4306

Query Match 100.0%; Score 86; DB 24; Length 134;  
Best Local Similarity 100.0%; Pred. No. 7.5e-06;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FGGFTGARKSARKLANQ 17  
|||  
Db 88 FGGFTGARKSARKLANQ 104

RESULT 15

US-60-160-203-6127  
; Sequence 6127, Application US/60160203  
; GENERAL INFORMATION:  
; APPLICANT: BONAZZI, VIVIEN  
; TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,  
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS  
AND  
; TITLE OF INVENTION: USES THEREOF  
; FILE REFERENCE: CL000116  
; CURRENT APPLICATION NUMBER: US/60/160,203  
; CURRENT FILING DATE: 1999-10-19  
; NUMBER OF SEQ ID NOS: 6374  
; SOFTWARE: FastSEQ for Windows Version 4.0  
; SEQ ID NO 6127  
; LENGTH: 134  
; TYPE: PRT  
; ORGANISM: HUMAN  
US-60-160-203-6127

Query Match 100.0%; Score 86; DB 24; Length 134;  
Best Local Similarity 100.0%; Pred. No. 7.5e-06;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FGGFTGARKSARKLANQ 17  
|||  
Db 88 FGGFTGARKSARKLANQ 104

Search completed: November 16, 2001, 15:49:20  
Job time: 307 sec

## 09011797 Interference Results

SEQ ID NO: 1

## SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	932	100.0	932	14	US-09-011-797-1	Sequence 1, Appli
2	546.4	58.6	1209	50	US-60-172-373-1728	Sequence 1728, Ap
3	323.6	34.7	554	20	US-09-534-843-5886	Sequence 5886, Ap
4	310	33.3	745	1	PCT-US01-18569-1506	Sequence 1506, Ap
5	308.6	33.1	644	49	US-60-160-202-2110	Sequence 2110, Ap
6	308.6	33.1	644	49	US-60-160-203-2940	Sequence 2940, Ap
7	308.6	33.1	645	49	US-60-160-202-1762	Sequence 1762, Ap
8	308.6	33.1	645	49	US-60-160-203-2171	Sequence 2171, Ap
9	277	29.7	423	18	US-09-431-517-13593	Sequence 13593, A
10	270.8	29.1	522	20	US-09-534-843-5889	Sequence 5889, Ap
11	267.6	28.7	589	21	US-09-540-229-112193	Sequence 112193,
12	267.6	28.7	589	26	US-09-668-337-251	Sequence 251, App
13	267.6	28.7	589	49	US-60-168-197-252	Sequence 252, App
14	264.2	28.3	462	16	US-09-248-797-6959	Sequence 6959, Ap
15	263.2	28.2	521	20	US-09-534-843-5888	Sequence 5888, Ap
16	240.4	25.8	503	20	US-09-534-843-5887	Sequence 5887, Ap
17	226.8	24.3	456	19	US-09-528-409-49212	Sequence 49212, A
18	226.8	24.3	456	32	US-09-933-524-49212	Sequence 49212, A
19	202	21.7	327	29	US-09-758-467-119	Sequence 119, App
20	197.8	21.2	492	16	US-09-235-076-2514	Sequence 2514, Ap
21	197.8	21.2	492	16	US-09-248-797-10027	Sequence 10027, A
22	197.8	21.2	492	17	US-09-332-782-2514	Sequence 2514, Ap
23	197.8	21.2	492	29	US-09-737-223-2514	Sequence 2514, Ap
24	194.2	20.8	471	16	US-09-287-618-35257	Sequence 35257, A
25	193.6	20.8	406	19	US-09-528-409-49211	Sequence 49211, A
26	193.6	20.8	406	32	US-09-933-524-49211	Sequence 49211, A
27	186.6	20.0	386	17	US-09-359-067-45870	Sequence 45870, A
28	172.6	18.5	411	16	US-09-289-768-1144	Sequence 1144, Ap
c 29	169.4	18.2	674	49	US-60-162-356-32	Sequence 32, Appl
c 30	169.4	18.2	674	49	US-60-163-123-70	Sequence 70, Appl
c 31	169.4	18.2	674	49	US-60-163-232-93	Sequence 93, Appl
c 32	168.4	18.1	673	49	US-60-162-242-380	Sequence 380, App
c 33	168.4	18.1	673	49	US-60-162-243-290	Sequence 290, App
c 34	168.4	18.1	673	49	US-60-162-356-550	Sequence 550, App
35	150.4	16.1	327	8	US-08-401-791A-13192	Sequence 13192, A
36	150.4	16.1	327	8	US-08-401-791B-13192	Sequence 13192, A
37	149.2	16.0	235	18	US-09-440-302-505	Sequence 505, App
38	133.4	14.3	499	1	PCT-US99-18054-50	Sequence 50, Appl
39	133.4	14.3	499	16	US-09-244-694-50	Sequence 50, Appl
40	127.6	13.7	186	3	US-07-925-932-490	Sequence 490, App
41	109.4	11.7	240	21	US-09-540-766-52725	Sequence 52725, A
42	99	10.6	143	29	US-09-758-467-26	Sequence 26, Appl
43	95.2	10.2	245	16	US-09-250-152-2974	Sequence 2974, Ap
44	95.2	10.2	301	15	US-09-145-501-1321	Sequence 1321, Ap
45	95.2	10.2	301	39	US-60-069-691-1321	Sequence 1321, Ap

## SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
c 1	164.6	17.7	305	5	US-09-954-456-1045	Sequence 1045, Ap
c 2	37.8	4.1	504	6	US-60-325-448-456	Sequence 456, App
c 3	35.6	3.8	2938	5	US-09-710-481-3	Sequence 3, Appli
4	35	3.8	343	5	US-09-388-906A-13517	Sequence 13517, A
5	35	3.8	3096	5	US-09-685-791-9	Sequence 9, Appli
6	35	3.8	3949	5	US-09-685-791-7	Sequence 7, Appli
c 7	34.8	3.7	2107	5	US-09-981-353-49	Sequence 49, Appl
c 8	34.6	3.7	436	5	US-09-922-340-7747	Sequence 7747, Ap



c	9	34.6	3.7	684	5	US-09-815-242-4034	Sequence 4034, Ap
c	10	34.4	3.7	396	5	US-09-970-966-42	Sequence 42, Appl
	11	34.4	3.7	437	5	US-09-850-716A-74	Sequence 74, Appl
	12	34.2	3.7	480	5	US-09-922-340-11753	Sequence 11753, A
c	13	34	3.6	473	5	US-09-922-340-7749	Sequence 7749, Ap
c	14	34	3.6	475	5	US-09-922-340-7748	Sequence 7748, Ap
c	15	33.8	3.6	3648	5	US-09-800-187-5	Sequence 5, Appli
c	16	33.6	3.6	494	5	US-09-850-716A-54	Sequence 54, Appl
	17	33.4	3.6	316	5	US-09-388-906A-21091	Sequence 21091, A
	18	33.4	3.6	455	5	US-09-922-340-11897	Sequence 11897, A
c	19	33.2	3.6	211	5	US-09-388-906A-16059	Sequence 16059, A
c	20	33.2	3.6	300	5	US-09-388-906A-22896	Sequence 22896, A
	21	33.2	3.6	607	6	US-60-325-448-3446	Sequence 3446, Ap
c	22	33	3.5	624	5	US-09-961-619-17	Sequence 17, Appl
	23	33	3.5	720	6	US-60-325-448-4174	Sequence 4174, Ap
c	24	32.8	3.5	2365	5	US-09-978-189-131	Sequence 131, App
c	25	32.8	3.5	2365	5	US-09-978-192-131	Sequence 131, App
c	26	32.8	3.5	2365	5	US-09-978-697-131	Sequence 131, App
c	27	32.8	3.5	2365	5	US-09-978-824-131	Sequence 131, App
c	28	32.4	3.5	237	5	US-09-815-242-4157	Sequence 4157, Ap
	29	32.4	3.5	264	5	US-09-815-242-1204	Sequence 1204, Ap
c	30	32.4	3.5	425	5	US-09-834-975-451	Sequence 451, App
	31	32.4	3.5	6420	5	US-09-937-837-2	Sequence 2, Appli
c	32	32.2	3.5	401	5	US-09-388-906A-22533	Sequence 22533, A
	33	32.2	3.5	7869	5	US-09-954-456-1921	Sequence 1921, Ap
	34	32.2	3.5	48841	5	US-09-844-653-32	Sequence 32, Appl
c	35	32	3.4	1025	5	US-09-969-730-72	Sequence 72, Appl
	36	32	3.4	1065	5	US-09-969-730-105	Sequence 105, App
	37	31.8	3.4	2567	6	US-60-325-448-258	Sequence 258, App
	38	31.8	3.4	88421	5	US-09-976-059-1	Sequence 1, Appli
	39	31.6	3.4	308	5	US-09-922-340-7098	Sequence 7098, Ap
c	40	31.6	3.4	434	5	US-09-954-456-2098	Sequence 2098, Ap
	41	31.6	3.4	990	5	US-09-815-242-7714	Sequence 7714, Ap
	42	31.6	3.4	1827	5	US-09-778-927A-1	Sequence 1, Appli
	43	31.6	3.4	2182	5	US-09-778-927A-2	Sequence 2, Appli
c	44	31.4	3.4	611	5	US-09-871-161-393	Sequence 393, App
	45	31.4	3.4	2173	5	US-09-525-998-14	Sequence 14, Appl

SEQ ID NO: 2

# SUMMARIES

Result No.	Score	% Match	Query Length	DB	ID	Description
1	86	100.0	17	9	US-08-514-451-5	Sequence 5, Appli
2	86	100.0	17	9	US-08-553-058A-5	Sequence 5, Appli
3	86	100.0	17	12	US-08-868-355-1	Sequence 1, Appli
4	86	100.0	17	13	US-08-927-328-1	Sequence 1, Appli
5	86	100.0	17	14	US-09-011-797-2	Sequence 2, Appli
6	86	100.0	17	14	US-09-048-916-5	Sequence 5, Appli
7	86	100.0	17	14	US-09-048-916-5	Sequence 5, Appli
8	86	100.0	17	14	US-09-048-916B-5	Sequence 5, Appli
9	86	100.0	17	15	US-09-114-620-1	Sequence 1, Appli
10	86	100.0	17	15	US-09-170-919-5	Sequence 5, Appli
11	86	100.0	17	17	US-09-341-590-39	Sequence 39, Appl
12	86	100.0	17	20	US-09-657-276-919	Sequence 919, App
13	86	100.0	17	20	US-09-657-276-926	Sequence 926, App
14	86	100.0	134	24	US-60-160-202-4306	Sequence 4306, Ap
15	86	100.0	134	24	US-60-160-203-6127	Sequence 6127, Ap
16	86	100.0	139	24	US-60-160-203-5358	Sequence 5358, Ap
17	86	100.0	155	24	US-60-160-202-3958	Sequence 3958, Ap
18	86	100.0	188	1	PCT-US01-18569-3681	Sequence 3681, Ap
19	83	96.5	17	13	US-08-927-328-4	Sequence 4, Appli
20	83	96.5	17	13	US-08-927-328-6	Sequence 6, Appli
21	83	96.5	17	13	US-08-927-328-8	Sequence 8, Appli
22	81	94.2	17	9	US-08-514-451-6	Sequence 6, Appli
23	81	94.2	17	9	US-08-553-058A-6	Sequence 6, Appli
24	81	94.2	17	14	US-09-048-916-6	Sequence 6, Appli

25	81	94.2	17	14	US-09-048-916-6	Sequence 6, Appli
26	81	94.2	17	14	US-09-048-916B-6	Sequence 6, Appli
27	81	94.2	17	15	US-09-114-620-4	Sequence 4, Appli
28	81	94.2	17	15	US-09-170-919-6	Sequence 6, Appli
29	80	93.0	16	13	US-08-927-328-3	Sequence 3, Appli
30	80	93.0	17	13	US-08-927-328-7	Sequence 7, Appli
31	74	86.0	17	13	US-08-927-328-5	Sequence 5, Appli
32	68	79.1	15	13	US-08-927-328-20	Sequence 20, Appl
33	68	79.1	16	13	US-08-927-328-17	Sequence 17, Appl
34	64	74.4	17	13	US-08-927-328-13	Sequence 13, Appl
35	63	73.3	14	13	US-08-927-328-30	Sequence 30, Appl
36	62	72.1	15	13	US-08-927-328-12	Sequence 12, Appl
37	62	72.1	16	13	US-08-927-328-11	Sequence 11, Appl
38	62	72.1	16	13	US-08-927-328-15	Sequence 15, Appl
39	61	70.9	16	13	US-08-927-328-19	Sequence 19, Appl
40	57	66.3	15	13	US-08-927-328-9	Sequence 9, Appli
41	57	66.3	15	13	US-08-927-328-14	Sequence 14, Appl
42	57	66.3	16	13	US-08-927-328-10	Sequence 10, Appl
43	57	66.3	16	13	US-08-927-328-16	Sequence 16, Appl
44	57	66.3	16	13	US-08-927-328-18	Sequence 18, Appl
45	44.5	51.7	319	15	US-09-198-452A-173	Sequence 173, App

# SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	86	100.0	17	5	US-09-929-986-1	Sequence 1, Appli
2	39	45.3	232	5	US-09-897-516-8334	Sequence 8334, Ap
3	38	44.2	225	5	US-09-897-516-4492	Sequence 4492, Ap
4	38	44.2	296	5	US-09-815-242-13677	Sequence 13677, A
5	38	44.2	454	5	US-09-897-516-7417	Sequence 7417, Ap
6	37	43.0	256	5	US-09-815-242-11383	Sequence 11383, A
7	37	43.0	416	5	US-09-897-516-4952	Sequence 4952, Ap
8	36	41.9	166	5	US-09-815-242-5181	Sequence 5181, Ap
9	36	41.9	566	5	US-09-897-516-6975	Sequence 6975, Ap
10	36	41.9	933	5	US-09-815-242-11817	Sequence 11817, A
11	36	41.9	1300	5	US-09-896-923-3	Sequence 3, Appli
12	35	40.7	230	5	US-09-815-242-10652	Sequence 10652, A
13	35	40.7	266	5	US-09-815-242-5346	Sequence 5346, Ap
14	35	40.7	266	5	US-09-815-242-12340	Sequence 12340, A
15	35	40.7	266	5	US-09-815-242-12989	Sequence 12989, A
16	34	39.5	87	5	US-09-815-242-11763	Sequence 11763, A
17	34	39.5	119	5	US-09-897-516-7459	Sequence 7459, Ap
18	34	39.5	163	5	US-09-615-846A-16	Sequence 16, Appl
19	34	39.5	332	5	US-09-815-242-10156	Sequence 10156, A
20	34	39.5	398	5	US-09-815-242-10390	Sequence 10390, A
21	34	39.5	398	5	US-09-897-516-6927	Sequence 6927, Ap
22	34	39.5	633	5	US-09-897-516-5052	Sequence 5052, Ap
23	34	39.5	741	1	PCT-US01-20545-13	Sequence 13, Appl
24	34	39.5	763	5	US-09-815-242-13643	Sequence 13643, A
25	33.5	39.0	99	5	US-09-795-668-35	Sequence 35, Appl
26	33.5	39.0	118	5	US-09-815-242-13222	Sequence 13222, A
27	33.5	39.0	125	5	US-09-795-668-30	Sequence 30, Appl
28	33.5	39.0	127	1	PCT-US01-31269-10	Sequence 10, Appl
29	33.5	39.0	226	5	US-09-954-314-4	Sequence 4, Appli
30	33.5	39.0	344	5	US-09-795-668-38	Sequence 38, Appl
31	33.5	39.0	456	5	US-09-795-668-17	Sequence 17, Appl
32	33.5	39.0	632	5	US-09-795-668-16	Sequence 16, Appl
33	33	38.4	130	5	US-09-815-242-5552	Sequence 5552, Ap
34	33	38.4	131	5	US-09-815-242-11426	Sequence 11426, A
35	33	38.4	132	5	US-09-815-242-12238	Sequence 12238, A
36	33	38.4	208	5	US-09-815-242-11425	Sequence 11425, A
37	33	38.4	280	5	US-09-815-242-5745	Sequence 5745, Ap
38	33	38.4	283	5	US-09-815-242-11268	Sequence 11268, A
39	33	38.4	296	5	US-09-815-242-13353	Sequence 13353, A
40	33	38.4	344	5	US-09-646-673A-177	Sequence 177, App
41	33	38.4	365	5	US-09-815-242-5593	Sequence 5593, Ap
42	33	38.4	365	5	US-09-815-242-12414	Sequence 12414, A

43	33	38.4	365	5	US-09-815-242-13025	Sequence 13025, A
44	33	38.4	670	5	US-09-468-646A-29	Sequence 29, Appl
45	33	38.4	702	5	US-09-897-516-7546	Sequence 7546, Ap

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#### SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	82	100.0	17	14	US-09-011-797-3	Sequence 3, Appli
2	82	100.0	17	15	US-09-114-620-2 <i>AW</i>	Sequence 2, Appli
3	82	100.0	134	24	US-60-160-202-4306	Sequence 4306, Ap
4	82	100.0	134	24	US-60-160-203-6127	Sequence 6127, Ap
5	82	100.0	139	24	US-60-160-203-5358	Sequence 5358, Ap
6	82	100.0	155	24	US-60-160-202-3958	Sequence 3958, Ap
7	76	92.7	188	1	PCT-US01-18569-3681	Sequence 3681, Ap
8	42	51.2	346	24	US-60-324-631-1830	Sequence 1830, Ap
9	40.5	49.4	2059	24	US-60-173-464-29693	Sequence 29693, A
10	40.5	49.4	2060	24	US-60-191-637-38941	Sequence 38941, A
11	40.5	49.4	2060	24	US-60-191-681-30175	Sequence 30175, A
12	39	47.6	62	23	US-09-950-083-3580	Sequence 3580, Ap
13	39	47.6	63	1	PCT-US00-06828-84	Sequence 84, Appl
14	39	47.6	159	16	US-09-270-767-41900	Sequence 41900, A
15	38.5	47.0	160	21	US-09-733-089-14467	Sequence 14467, A
16	38.5	47.0	160	22	US-09-816-660-14467	Sequence 14467, A
17	38	46.3	138	1	PCT-US01-08631-38669	Sequence 38669, A
18	38	46.3	236	19	US-09-540-236-3493	Sequence 3493, Ap
19	38	46.3	236	24	US-60-128-476-4429	Sequence 4429, Ap
20	38	46.3	327	16	US-09-248-796-16639	Sequence 16639, A
21	38	46.3	341	1	PCT-US01-08631-38673	Sequence 38673, A
22	38	46.3	379	1	PCT-US01-08631-50988	Sequence 50988, A
23	38	46.3	384	1	PCT-US01-08631-38671	Sequence 38671, A
24	38	46.3	414	1	PCT-US01-16450-2318	Sequence 2318, Ap
25	38	46.3	414	1	PCT-US01-16450A-2318	Sequence 2318, Ap
26	38	46.3	441	18	US-09-489-039A-10210	Sequence 10210, A
27	38	46.3	494	17	US-09-328-352-5284	Sequence 5284, Ap
28	38	46.3	497	22	US-09-805-020-37	Sequence 37, Appl
29	38	46.3	543	1	PCT-US01-08631-53304	Sequence 53304, A
30	38	46.3	731	18	US-09-494-810-7	Sequence 7, Appli
31	38	46.3	731	18	US-09-494-810A-7	Sequence 7, Appli
32	38	46.3	741	22	US-09-833-790-432	Sequence 432, App
33	38	46.3	757	1	PCT-US01-08631-38674	Sequence 38674, A
34	37	45.1	24	21	US-09-724-059-413838	Sequence 413838,
35	37	45.1	24	21	US-09-724-059-414216	Sequence 414216,
36	37	45.1	24	21	US-09-724-059-421715	Sequence 421715,
37	37	45.1	24	21	US-09-724-059-421967	Sequence 421967,
38	37	45.1	24	21	US-09-724-059-446796	Sequence 446796,
39	37	45.1	24	21	US-09-724-059-449316	Sequence 449316,
40	37	45.1	64	16	US-09-248-796-27227	Sequence 27227, A
41	37	45.1	70	24	US-60-186-662-622	Sequence 622, App
42	37	45.1	97	1	PCT-US01-14827-12232	Sequence 12232, A
43	37	45.1	119	18	US-09-474-434-1050	Sequence 1050, Ap
44	37	45.1	119	24	US-60-173-686-1050	Sequence 1050, Ap
45	37	45.1	123	15	US-09-134-001C-3630	Sequence 3630, Ap

#### SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	36	43.9	102	5	US-09-815-242-5417	Sequence 5417, Ap
2	36	43.9	102	5	US-09-815-242-12559	Sequence 12559, A
3	36	43.9	102	5	US-09-815-242-12960	Sequence 12960, A
4	35	42.7	375	5	US-09-815-242-11878	Sequence 11878, A

5	34	41.5	129	5	US-09-461-436-327	Sequence 327, App
6	34	41.5	362	5	US-09-461-436-374	Sequence 374, App
7	34	41.5	362	5	US-09-779-679-28	Sequence 28, Appl
8	34	41.5	555	5	US-09-978-189-109	Sequence 109, App
9	34	41.5	555	5	US-09-978-192-109	Sequence 109, App
10	34	41.5	555	5	US-09-978-697-109	Sequence 109, App
11	34	41.5	555	5	US-09-978-824-109	Sequence 109, App
12	33	40.2	315	5	US-09-897-516-5136	Sequence 5136, Ap
13	33	40.2	371	5	US-09-897-516-5976	Sequence 5976, Ap
14	33	40.2	578	5	US-09-815-242-12501	Sequence 12501, A
15	33	40.2	689	5	US-09-897-516-6629	Sequence 6629, Ap
16	32	39.0	376	5	US-09-897-516-4587	Sequence 4587, Ap
17	32	39.0	439	5	US-09-815-242-13917	Sequence 13917, A
18	32	39.0	1905	5	US-09-897-516-6551	Sequence 6551, Ap
19	31	37.8	439	5	US-09-815-242-10063	Sequence 10063, A
20	31	37.8	465	5	US-09-897-516-6575	Sequence 6575, Ap
21	31	37.8	497	5	US-09-981-353-105	Sequence 105, App
22	31	37.8	651	5	US-09-897-516-7732	Sequence 7732, Ap
23	31	37.8	657	5	US-09-815-242-13436	Sequence 13436, A
24	31	37.8	686	5	US-09-897-516-6833	Sequence 6833, Ap
25	31	37.8	975	5	US-09-897-516-7681	Sequence 7681, Ap
26	31	37.8	1084	5	US-09-800-187-2	Sequence 2, Appli
27	30	36.6	142	5	US-09-897-516-6053	Sequence 6053, Ap
28	30	36.6	180	5	US-09-815-242-5657	Sequence 5657, Ap
29	30	36.6	185	5	US-09-815-242-12269	Sequence 12269, A
30	30	36.6	224	5	US-09-897-516-8168	Sequence 8168, Ap
31	30	36.6	271	5	US-09-897-516-4443	Sequence 4443, Ap
32	30	36.6	290	5	US-09-815-242-13873	Sequence 13873, A
33	30	36.6	362	5	US-09-897-516-6719	Sequence 6719, Ap
34	30	36.6	441	5	US-09-897-516-7789	Sequence 7789, Ap
35	30	36.6	446	5	US-09-897-516-7790	Sequence 7790, Ap
36	30	36.6	531	5	US-09-897-516-7946	Sequence 7946, Ap
37	30	36.6	652	5	US-09-815-242-13317	Sequence 13317, A
38	30	36.6	652	5	US-09-815-242-13673	Sequence 13673, A
39	30	36.6	777	5	US-09-815-242-4894	Sequence 4894, Ap
40	30	36.6	871	5	US-09-815-242-5274	Sequence 5274, Ap
41	30	36.6	872	5	US-09-815-242-12333	Sequence 12333, A
42	30	36.6	914	5	US-09-815-242-10897	Sequence 10897, A
43	30	36.6	1089	5	US-09-769-987-2	Sequence 2, Appli
44	29.5	36.0	310	5	US-09-886-055-153	Sequence 153, App
45	29.5	36.0	610	5	US-09-897-516-8116	Sequence 8116, Ap

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# SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	44	100.0	8	14	US-09-011-797-4	Sequence 4, Appli
2	44	100.0	8	15	US-09-114-620-3	Sequence 3, Appli
3	44	100.0	134	24	US-60-160-202-4306	Sequence 4306, Ap
4	44	100.0	134	24	US-60-160-203-6127	Sequence 6127, Ap
5	44	100.0	139	24	US-60-160-203-5358	Sequence 5358, Ap
6	44	100.0	155	24	US-60-160-202-3958	Sequence 3958, Ap
7	35	79.5	170	16	US-09-248-796-15906	Sequence 15906, A
8	35	79.5	377	16	US-09-270-767-34332	Sequence 34332, A
9	35	79.5	377	16	US-09-270-767-49549	Sequence 49549, A
10	35	79.5	377	16	US-09-270-849B-180627	Sequence 180627,
11	35	79.5	559	24	US-60-248-505-822	Sequence 822, App
12	35	79.5	601	24	US-60-248-505-1284	Sequence 1284, Ap
13	35	79.5	635	16	US-09-252-991A-31646	Sequence 31646, A
14	35	79.5	1426	24	US-60-191-637-40785	Sequence 40785, A
15	35	79.5	1426	24	US-60-191-700-720	Sequence 720, App
16	35	79.5	1542	24	US-60-167-324-686	Sequence 686, App
17	35	79.5	1542	24	US-60-173-386-656	Sequence 656, App
18	35	79.5	1542	24	US-60-175-871-737	Sequence 737, App
19	35	79.5	1542	24	US-60-184-775-671	Sequence 671, App
20	34	77.3	99	1	PCT-US01-08631-50558	Sequence 50558, A

21	34	77.3	152	16	US-09-270-767-38731	Sequence 38731, A
22	34	77.3	152	16	US-09-270-767-53948	Sequence 53948, A
23	34	77.3	152	16	US-09-270-849B-189754	Sequence 189754, A
24	34	77.3	156	1	PCT-US01-08631-50559	Sequence 50559, A
25	34	77.3	224	15	US-09-149-476-370	Sequence 370, App
26	34	77.3	224	22	US-09-809-391-370	Sequence 370, App
27	34	77.3	224	22	US-09-882-171-370	Sequence 370, App
28	34	77.3	230	22	US-09-855-768-673	Sequence 673, App
29	34	77.3	349	24	US-60-191-637-30778	Sequence 30778, A
30	34	77.3	349	24	US-60-191-681-24517	Sequence 24517, A
31	34	77.3	450	24	US-60-173-464-24182	Sequence 24182, A
32	34	77.3	451	24	US-60-175-691-147	Sequence 147, App
33	34	77.3	460	24	US-60-215-161-7857	Sequence 7857, Ap
34	34	77.3	492	20	US-09-618-893-147	Sequence 147, App
35	34	77.3	584	1	PCT-US01-08631-50562	Sequence 50562, A
36	34	77.3	638	1	PCT-US01-08631-56551	Sequence 56551, A
37	33	75.0	120	16	US-09-248-796-17350	Sequence 17350, A
38	33	75.0	210	17	US-09-328-352-4351	Sequence 4351, Ap
39	33	75.0	251	13	US-08-993-002A-8279	Sequence 8279, Ap
40	33	75.0	260	1	PCT-US01-19110-1039	Sequence 1039, Ap
41	33	75.0	260	22	US-09-880-748-1039	Sequence 1039, Ap
42	33	75.0	279	10	US-08-625-811-1714	Sequence 1714, Ap
43	33	75.0	279	13	US-08-993-002A-8280	Sequence 8280, Ap
44	33	75.0	306	16	US-09-252-691-6233	Sequence 6233, Ap
45	33	75.0	306	16	US-09-252-691C-6233	Sequence 6233, Ap

# SUMMARIES

Result No.	Score	% Match	Query Length	DB	ID	Description
1	34	77.3	460	5	US-09-897-516-7857	Sequence 7857, Ap
2	33	75.0	1395	5	US-09-800-198-56	Sequence 56, Appl
3	31	70.5	148	5	US-09-615-846A-17	Sequence 17, Appl
4	31	70.5	314	5	US-09-725-945-6	Sequence 6, Appli
5	31	70.5	1041	5	US-09-978-189-498	Sequence 498, App
6	31	70.5	1041	5	US-09-978-192-498	Sequence 498, App
7	31	70.5	1041	5	US-09-978-697-498	Sequence 498, App
8	31	70.5	1041	5	US-09-978-824-498	Sequence 498, App
9	30	68.2	745	5	US-09-897-516-7799	Sequence 7799, Ap
10	30	68.2	934	5	US-09-971-490-2	Sequence 2, Appli
11	29	65.9	212	5	US-09-897-516-7312	Sequence 7312, Ap
12	29	65.9	355	5	US-09-609-360C-26	Sequence 26, Appl
13	29	65.9	355	5	US-09-345-473E-26	Sequence 26, Appl
14	29	65.9	554	5	US-09-815-242-12033	Sequence 12033, A
15	28	63.6	151	5	US-09-815-242-11515	Sequence 11515, A
16	28	63.6	200	5	US-09-545-199D-63	Sequence 63, Appl
17	28	63.6	222	5	US-09-897-516-5345	Sequence 5345, Ap
18	28	63.6	242	5	US-09-815-242-10974	Sequence 10974, A
19	28	63.6	244	5	US-09-815-242-10126	Sequence 10126, A
20	28	63.6	334	5	US-09-897-516-6895	Sequence 6895, Ap
21	28	63.6	440	5	US-09-815-242-5131	Sequence 5131, Ap
22	28	63.6	581	5	US-09-897-516-6038	Sequence 6038, Ap
23	28	63.6	613	5	US-09-897-516-5750	Sequence 5750, Ap
24	28	63.6	639	5	US-09-815-242-5390	Sequence 5390, Ap
25	28	63.6	646	5	US-09-815-242-12304	Sequence 12304, A
26	28	63.6	722	5	US-09-897-516-6067	Sequence 6067, Ap
27	28	63.6	734	5	US-09-545-199D-117	Sequence 117, App
28	28	63.6	1114	5	US-09-840-743-14	Sequence 14, Appl
29	27	61.4	75	5	US-09-969-730-139	Sequence 139, App
30	27	61.4	112	4	US-08-859-648-19	Sequence 19, Appl
31	27	61.4	112	4	US-08-859-648-25	Sequence 25, Appl
32	27	61.4	112	4	US-08-859-648-29	Sequence 29, Appl
33	27	61.4	112	4	US-08-859-648-33	Sequence 33, Appl
34	27	61.4	177	5	US-09-938-497-15	Sequence 15, Appl
35	27	61.4	218	5	US-09-897-516-5887	Sequence 5887, Ap
36	27	61.4	238	4	US-08-721-612C-19	Sequence 19, Appl
37	27	61.4	276	5	US-09-897-516-4756	Sequence 4756, Ap
38	27	61.4	277	5	US-09-897-516-7437	Sequence 7437, Ap

39	27	61.4	324	5	US-09-815-242-11345	Sequence 11345, A
40	27	61.4	330	5	US-09-815-242-5293	Sequence 5293, Ap
41	27	61.4	333	5	US-09-815-242-12121	Sequence 12121, A
42	27	61.4	370	5	US-09-815-242-5528	Sequence 5528, Ap
43	27	61.4	370	5	US-09-815-242-12353	Sequence 12353, A
44	27	61.4	476	5	US-09-960-643-2	Sequence 2, Appli
45	27	61.4	636	5	US-09-815-242-5838	Sequence 5838, Ap